SA Code

nohup IntaRNA -t SA\_1.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_1\_targets.txt &> nohuphsa-SA\_1.out &

nohup IntaRNA -t SA\_2.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_2\_targets.txt &> nohuphsa-SA\_2.out &

nohup IntaRNA -t SA\_3.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_3\_targets.txt &> nohuphsa-SA\_3.out &

nohup IntaRNA -t SA\_4.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_4\_targets.txt &> nohuphsa-SA\_4.out &

nohup IntaRNA -t SA\_5.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_5\_targets.txt &> nohuphsa-SA\_5.out &

nohup IntaRNA -t SA\_6.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_6\_targets.txt &> nohuphsa-SA\_6.out &

nohup IntaRNA -t SA\_7.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_7\_targets.txt &> nohuphsa-SA\_7.out &

nohup IntaRNA -t SA\_8.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_8\_targets.txt &> nohuphsa-SA\_8.out &

nohup IntaRNA -t SA\_9.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_9\_targets.txt &> nohuphsa-SA\_9.out &

nohup IntaRNA -t SA\_10.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_10\_targets.txt &> nohuphsa-SA\_10.out &

nohup IntaRNA -t SA\_11.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_11\_targets.txt &> nohuphsa-SA\_11.out &

nohup IntaRNA -t SA\_12.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SA\_12\_targets.txt &> nohuphsa-SA\_12.out &